

Fitting von Bertalanffy Growth Models with Fixed Sex Effects

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Load Required Packages and Data

```
LMBL <- read.csv("Data/Clean-Data/2016_largemouth-bass_long-format.csv") %>%
  select(FID:BI.len) %>%
  arrange(FID,Agei)

### Making factors factors
LMBL$FID <- factor(LMBL$FID)
LMBL$Site <- factor(LMBL$Site)
LMBL$SEXCON <- factor(LMBL$SEXCON)
LMBL$Sex <- factor(LMBL$Sex)

str(LMBL)

## 'data.frame': 337 obs. of 11 variables:
## $ FID : Factor w/ 126 levels "1","2","3","4",...: 1 1 1 1 2 2 2 2 3 3 ...
## $ Site : Factor w/ 11 levels "2","4","6","8",...: 6 6 6 6 6 6 6 6 6 6 ...
## $ AgeCap: int 4 4 4 4 4 4 4 4 4 4 ...
## $ RadCap: num 0.94 0.94 0.94 0.94 0.988 ...
## $ LenCap: int 347 347 347 347 292 292 292 292 348 348 ...
## $ WTg : int 658 658 658 658 415 415 415 415 557 557 ...
## $ SEXCON: Factor w/ 5 levels "0","1","3","6",...: 5 5 5 5 3 3 3 3 3 3 ...
## $ Sex : Factor w/ 3 levels "0","1","2": 3 3 3 3 2 2 2 2 2 2 ...
## $ Agei : int 1 2 3 4 1 2 3 4 1 2 ...
## $ Radi : num 0.433 0.69 0.803 0.927 0.567 ...
## $ BI.len: num 155 252 295 342 165 ...

headtail(LMBL)

## FID Site AgeCap RadCap LenCap WTg SEXCON Sex Agei Radi BI.len
## 1 1 11 4 0.9402 347 658 8 2 1 0.4328 154.6790
## 2 1 11 4 0.9402 347 658 8 2 2 0.6898 252.0903
## 3 1 11 4 0.9402 347 658 8 2 3 0.8028 294.9210
## 335 132 15972 7 1.0474 395 971 3 1 5 0.9567 359.9617
## 336 132 15972 7 1.0474 395 971 3 1 6 1.0119 381.2860
## 337 132 15972 7 1.0474 395 971 3 1 7 1.0365 390.7892
```

Creating the von Bertalanffy function.

```
LVB <- function(x, Linf, K, t0){
  y = Linf * (1 - exp(-K * (x - t0)))
  y
}
LVB <- vbFuns()

LVB(5, 422.8, 0.39, -0.40)
```

```
## [1] 371.3351
LVB(5, Linf = c(422.8, 0.39, -0.40)) ### Should be the same output

## [1] 371.3351
```

Modeling Fixed Effects for Sex

Lets look at the von Bertalanffy growth model fits with a fixed sex term on the parameter estimates.

In order to fit the sex model I will have to remove the individual with no sex. In order to compare the sex model to the no sex model I will have to re-fit the no sex model without the individual with unknown sex. This means making new df excluding this individual. Then, making a new grouped data object. Finnaly, I will need to rerun the nlme function with out this fish to get the nlme.mod2 output [*This was All done in the fit-von-bertalanffy-growth-model.rmd file*].

Removing the individual with no sex (FID=89??)

```
### Just looking at data
head(LMBL)

##   FID Site AgeCap RadCap LenCap WTg SEXCON Sex Agei  Radi  BI.len
## 1   1   11     4 0.9402   347 658     8   2    1 0.4328 154.6790
## 2   1   11     4 0.9402   347 658     8   2    2 0.6898 252.0903
## 3   1   11     4 0.9402   347 658     8   2    3 0.8028 294.9210
## 4   1   11     4 0.9402   347 658     8   2    4 0.9269 341.9589
## 5   2   11     4 0.9884   292 415     3   1    1 0.5665 164.5096
## 6   2   11     4 0.9884   292 415     3   1    2 0.7181 210.3203

### Finding fish with unknown sex
(unknown.sex <- filterD(LMBL, Sex==0))

##   FID Site AgeCap RadCap LenCap WTg SEXCON Sex Agei  Radi  BI.len
## 1  89    8     1 0.4328   136 38      0   0    1 0.3983 124.5434

### Getting row number for fish with the unknown sex
(FID89 <- as.numeric(row.names(LMBL[LMBL$Sex==0,])))

## [1] 230

### removing the fish with unknown sex from the data set
length(LMBL$FID) ### just seeing the number of rows in the data set

## [1] 337

length(unique(LMBL$FID)) ### just seeing the number of fish

## [1] 126

LMBL <- LMBL[-c(FID89),]
length(LMBL$FID)

## [1] 336

length(unique(LMBL$FID)) ### Good! looks like only FID 89 was removed

## [1] 125
```

```
### Lets make sure there is no empty row in my data
LMBL <- filterD(LMBL,!is.na(FID))
### and lets just take a quick look at the data
str(LMBL)
```

```
## 'data.frame': 336 obs. of 11 variables:
## $ FID : Factor w/ 125 levels "1","2","3","4",...: 1 1 1 1 2 2 2 2 3 3 ...
## $ Site : Factor w/ 11 levels "2","4","6","8",...: 6 6 6 6 6 6 6 6 6 6 ...
## $ AgeCap: int 4 4 4 4 4 4 4 4 4 4 ...
## $ RadCap: num 0.94 0.94 0.94 0.94 0.988 ...
## $ LenCap: int 347 347 347 347 292 292 292 292 348 348 ...
## $ WTg : int 658 658 658 658 415 415 415 415 557 557 ...
## $ SEXCON: Factor w/ 4 levels "1","3","6","8": 4 4 4 4 2 2 2 2 2 2 ...
## $ Sex : Factor w/ 2 levels "1","2": 2 2 2 2 1 1 1 1 1 1 ...
## $ Agei : int 1 2 3 4 1 2 3 4 1 2 ...
## $ Radi : num 0.433 0.69 0.803 0.927 0.567 ...
## $ BI.len: num 155 252 295 342 165 ...
```

```
headtail(LMBL)
```

```
##      FID Site AgeCap RadCap LenCap WTg SEXCON Sex Agei Radi BI.len
## 1      1  11      4 0.9402   347 658      8  2    1 0.4328 154.6790
## 2      1  11      4 0.9402   347 658      8  2    2 0.6898 252.0903
## 3      1  11      4 0.9402   347 658      8  2    3 0.8028 294.9210
## 334 132 15972      7 1.0474   395 971      3  1    5 0.9567 359.9617
## 335 132 15972      7 1.0474   395 971      3  1    6 1.0119 381.2860
## 336 132 15972      7 1.0474   395 971      3  1    7 1.0365 390.7892
```

Remake Grouped Data Object

```
datgr = groupedData(BI.len ~ Agei|FID, data = LMBL,
                    labels = list(x = "Age", y = "Size"),
                    units = list(x = "(Years)", y = "(mm)"))
```

Full Sex Model, {sexmod.lkt}

Sex terms on all model parameters

I'm going to skip this for now it seems to be taking forever. I have this model output already converged, however, I did so with ML estimation instead of REML so I cannot compare with other model fits. I don't think this model should be used anyways since I'm not aware of a biological reason male and female largemouth bass would have a different t_0 .

```
sexmod.lkt <- nlme::nlme(BI.len ~ LVB(Agei, Linf, K, t0), data = datgr,
                        fixed = list(Linf ~ Sex-1, K ~ Sex-1, t0 ~ Sex-1),
                        random = Linf+K+t0 ~ 1,
                        start = list(fixed =
                                      c(Linf = c(389.3647,389.3647),
                                        K = c(0.4359,0.4359),
                                        t0 = c(-0.3127,-0.3127))),
                        method= "REML",
                        control=list(opt="nlsminb",
                                    maxIter=1000,
                                    pnlsMaxIter=100,
                                    msMaxIter=100,
```

```

niterEM=100))
save(sexmod.lkt,
      file = "model-output/sexmod.lkt.rda")

```

$\{L_\infty, K\}$ Sex Model, {sexmod.lk}

I'm going to skip this for now it seems to be taking forever. I have this model output already converged, however, I did so with ML estimation instead of REML so I cannot compare with other model fits.

Unfortunately, I do think this is the sex model that would make the most sense biologically except for maybe the L_∞ of K sex models.

```

sexmod.lk <- nlme::nlme(BI.len ~ LVB(Agei, Linf, K, t0), data = datgr,
                      fixed = list(Linf ~ Sex-1, K ~ Sex-1, t0 ~ 1),
                      random = Linf+K+t0 ~ 1,
                      start = list(fixed =
                                c(Linf = c(389.3647,389.3647),
                                  K = c(0.4359,0.4359),
                                  t0 = -0.3127)),
                      method= "REML",
                      control=list(opt="nlminb",
                                   maxIter=1000,
                                   pnlsMaxIter=100,
                                   msMaxIter=100,
                                   niterEM=100))
save(sexmod.lk,
      file = "model-output/sexmod.lk.rda")

```

$\{L_\infty, t_0\}$ Sex Model, {sexmod.lt}

```

sexmod.lt <- nlme::nlme(BI.len ~ LVB(Agei, Linf, K, t0), data = datgr,
                      fixed = list(Linf ~ Sex-1, K ~ 1, t0 ~ Sex-1),
                      random = Linf+K+t0 ~ 1,
                      start = list(fixed =
                                c(Linf = c(389.3647,389.3647),
                                  K = 0.4359,
                                  t0 = c(-0.3127,-0.3127))),
                      method= "REML",
                      control=list(opt="nlminb",
                                   maxIter=1000,
                                   pnlsMaxIter=100,
                                   msMaxIter=100,
                                   niterEM=100))
#save(sexmod.lt,
#      file = "model-output/sexmod.lt.rda")

```

$\{K, t_0\}$, Sex Model, {sexmod.kt}

```

sexmod.kt <- nlme::nlme(BI.len ~ LVB(Agei, Linf, K, t0), data = datgr,
                      fixed = list(Linf ~ 1, K ~ Sex-1, t0 ~ Sex-1),

```

```

random = Linf+K+t0 ~ 1,
start = list(fixed =
              c(Linf = 389.3647,
                K = c(0.4359,0.4359),
                t0 = c(-0.3127,-0.3127))),
method= "REML",
control=list(opt="nlminb",
             maxIter=1000,
             pnlsMaxIter=100,
             msMaxIter=100,
             niterEM=100))

#save(sexmod.kt,
#      file = "model-output/sexmod.kt.rda")

```

$\{L_\infty\}$, Sex Model, {sexmod.l}

```

sexmod.l <- nlme::nlme(BI.len ~ LVB(Agei, Linf, K, t0), data = datgr,
                      fixed = list(Linf ~ Sex-1, K ~ 1, t0 ~ 1),
                      random = Linf+K+t0 ~ 1,
                      start = list(fixed =
                                    c(Linf = c(389.3647,389.3647),
                                      K = 0.4359,
                                      t0 = -0.3127)),
                      method= "REML",
                      control=list(opt="nlminb",
                                   maxIter=1000,
                                   pnlsMaxIter=100,
                                   msMaxIter=100,
                                   niterEM=100))

#save(sexmod.l,
#      file = "model-output/sexmod.l.rda")

```

$\{K\}$, Sex Model, {sexmod.k}

```

sexmod.k <- nlme::nlme(BI.len ~ LVB(Agei, Linf, K, t0), data = datgr,
                      fixed = list(Linf ~ 1, K ~ Sex-1, t0 ~ 1),
                      random = Linf+K+t0 ~ 1,
                      start = list(fixed =
                                    c(Linf = 389.3647,
                                      K = c(0.4359,0.4359),
                                      t0 = -0.3127)),
                      method= "REML",
                      control=list(opt="nlminb",
                                   maxIter=1000,
                                   pnlsMaxIter=100,
                                   msMaxIter=100,
                                   niterEM=100))

#save(sexmod.k,
#      file = "model-output/sexmod.k.rda")

```

$\{t_0\}$, Sex Model, {sexmod.t}

```
sexmod.t <- nlme::nlme(BI.len ~ LVB(Agei, Linf, K, t0), data = datgr,
  fixed = list(Linf ~ 1, K ~ 1, t0 ~ Sex-1),
  random = Linf+K+t0 ~ 1,
  start = list(fixed =
    c(Linf = 389.3647,
      K = c(0.4359, 0.4359),
      t0 = -0.3127)),
  method= "REML",
  control=list(opt="nlsminb",
    maxIter=1000,
    pnlsMaxIter=100,
    msMaxIter=100,
    niterEM=100))

#save(sexmod.t,
#      file = "model-output/sexmod.t.rda")
```

zero intercept sex models ???

I will now try fitting the same sex models with a zero intercept (I think this is right might be slope). this is done by removing the -1 from VBPARAM ~ Sex-1 argument.

Rather than go through all of these right now which maybe I should I will just pick my favorites and go from there.

$\{L_\infty, K\}$ Intercept Sex Model {sexmod.lk.int}

```
sexmod.lk.int <- nlme::nlme(BI.len ~ LVB(Agei, Linf, K, t0), data = datgr,
  fixed = list(Linf ~ Sex, K ~ Sex, t0 ~ 1),
  random = Linf+K+t0 ~ 1,
  start = list(fixed =
    c(Linf = c(389.3647, 389.3647),
      K = c(0.4359, 0.4359),
      t0 = -0.3127)),
  method= "REML",
  control=list(opt="nlsminb",
    maxIter=100,
    pnlsMaxIter=1000,
    msMaxIter=100,
    niterEM=100))

save(sexmod.lk.int,
  file = "model-output/sexmod.lk.int.rda")
```

$\{L_\infty\}$ Intercept Sex Model {sexmod.l.int}

```
sexmod.l.int <- nlme::nlme(BI.len ~ LVB(Agei, Linf, K, t0), data = datgr,
  fixed = list(Linf ~ Sex, K ~ 1, t0 ~ 1),
  random = Linf+K+t0 ~ 1,
  start = list(fixed =
```

```

c(Linf = c(389.3647,389.3647),
  K = 0.4359,
  t0 = -0.3127)),
method= "REML",
control=list(opt="nlminb",
             maxIter=100,
             pnlsMaxIter=100,
             msMaxIter=100,
             niterEM=100))

#save(sexmod.l.int,
#      file = "model-output/sexmod.l.int.rda")

```

{K} Intercept Sex Model {sexmod.k.int}

```

sexmod.k.int <- nlme::nlme(BI.len ~ LVB(Agei, Linf, K, t0), data = datgr,
  fixed = list(Linf ~ 1, K ~ Sex, t0 ~ 1),
  random = Linf+K+t0 ~ 1,
  start = list(fixed =
    c(Linf = 389.3647,
      K = c(0.4359,0.4359),
      t0 = -0.3127)),
  method= "REML",
  control=list(opt="nlminb",
              maxIter=100,
              pnlsMaxIter=1000,
              msMaxIter=100,
              niterEM=100))

save(sexmod.k.int,
      file = "model-output/sexmod.k.int.rda")

```

Look at Model Fits

```

load("model-output/nlme.mod2.rda")

#load("model-output/sexmod.lkt.rda")
#load("model-output/sexmod.lk.rda")
load("model-output/sexmod.lt.rda")
load("model-output/sexmod.kt.rda")
load("model-output/sexmod.l.rda")
load("model-output/sexmod.k.rda")
load("model-output/sexmod.t.rda")

#load("model-output/sexmod.lkt.int.rda")
#load("model-output/sexmod.lk.int.rda")
#load("model-output/sexmod.lt.int.rda")
#load("model-output/sexmod.kt.int.rda")
load("model-output/sexmod.l.int.rda")
#load("model-output/sexmod.k.int.rda")
#load("model-output/sexmod.t.int.rda")

```

```
## [1] "Iterations = 11"
```

```

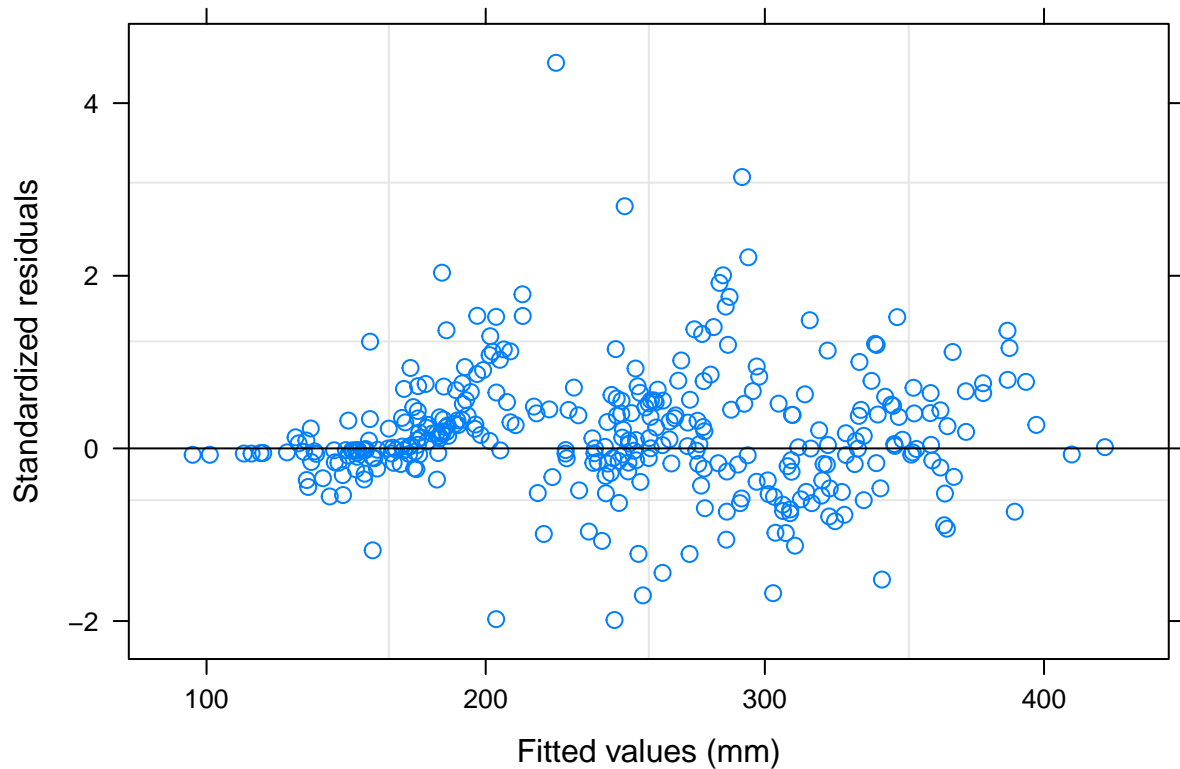
## Nonlinear mixed-effects model fit by REML
##   Model: BI.len ~ LVB(Agei, Linf, K, t0)
##   Data: datgr
##           AIC      BIC    logLik
##   2809.756 2851.613 -1393.878
##
## Random effects:
##   Formula: list(Linf ~ 1, K ~ 1, t0 ~ 1)
##   Level: FID
##   Structure: General positive-definite, Log-Cholesky parametrization
##           StdDev      Corr
## Linf.(Intercept) 60.7965255 Ln.(I) K
## K                 0.1283636 -0.880
## t0                0.4334910 -0.675  0.857
## Residual         4.7920294
##
## Fixed effects: list(Linf ~ Sex - 1, K ~ 1, t0 ~ 1)
##           Value Std.Error DF  t-value p-value
## Linf.Sex1 425.4479  8.432956 208 50.45062    0
## Linf.Sex2 439.0623  8.546468 208 51.37354    0
## K          0.3855  0.015426 208 24.98993    0
## t0        -0.3887  0.043652 208 -8.90441    0
##   Correlation:
##           Lnf.S1 Lnf.S2 K
## Linf.Sex2  0.695
## K          -0.824 -0.810
## t0         -0.582 -0.543  0.811
##
## Standardized Within-Group Residuals:
##           Min      Q1      Med      Q3      Max
## -1.98806932 -0.16716238  0.08612533  0.49651811  4.46707175
##
## Number of Observations: 336
## Number of Groups: 125
##
## Approximate 95% confidence intervals
##
##   Fixed effects:
##           lower      est.      upper
## Linf.Sex1 408.8228527 425.4478748 442.0728969
## Linf.Sex2 422.2135142 439.0623179 455.9111216
## K          0.3550875  0.3854992  0.4159109
## t0        -0.4747482 -0.3886919 -0.3026356
## attr("label")
## [1] "Fixed effects:"
##
##   Random Effects:
##   Level: FID
##           lower      est.      upper
## sd(Linf.(Intercept)) 47.29589478 60.7965255 78.1509163
## sd(K)                0.08225568  0.1283636  0.2003170
## sd(t0)               0.36854094  0.4334910  0.5098876
## cor(Linf.(Intercept),K) -0.95392287 -0.8799812 -0.7052801
## cor(Linf.(Intercept),t0) -0.85453505 -0.6745742 -0.3500078

```



```
## cor(K,t0)                0.71194480  0.8565798  0.9314982
##
## Within-group standard error:
##   lower    est.    upper
## 3.978896 4.792029 5.771337

##      numDF denDF   F-value p-value
## Linf.Sex    2   208 20992.203 <.0001
## K            1   208 3025.994 <.0001
## t0           1   208   79.289 <.0001
```



```
fixef(sexmod.1)
```

```
##   Linf.Sex1  Linf.Sex2      K      t0
## 425.4478748 439.0623179  0.3854992 -0.3886919
```

```
ranef(sexmod.1)
```

```
##      Linf.(Intercept)      K      t0
## 88      -96.09305881  0.2496292093  0.8778496208
## 87      -89.71634582  0.2299266925  0.8060792419
## 85      -82.95964757  0.2115588716  0.7468871005
## 86      -77.70709952  0.1958531873  0.6897164706
## 83      -78.14569333  0.1979820667  0.7007736752
## 27      -77.33178413  0.1957044398  0.6930647115
## 69      -69.82363331  0.1749195555  0.6230373032
## 84      -64.00480662  0.1590706823  0.5699787840
## 82      -55.36020645  0.1352840741  0.4867700302
## 110     -41.59308625  0.0994054508  0.3675613527
## 76      -44.97292140  0.1085899502  0.4020755279
## 80      -44.23785084  0.1066780798  0.3957214618
```

## 74	-43.08478201	0.1036844814	0.3857683022
## 78	-42.36027990	0.1018068294	0.3795228400
## 77	-42.07756079	0.1010748338	0.3770872465
## 75	-39.61713785	0.0947213152	0.3559277853
## 72	-31.44858559	0.0735853195	0.2812061103
## 121	-38.36861410	0.0915087775	0.3452123299
## 73	-14.78535586	0.0326456096	0.1416824757
## 7	-9.33227805	0.0199076831	0.0919711559
## 70	2.59650888	-0.0046519742	-0.0287300445
## 35	9.05285476	-0.0159100796	-0.0958419431
## 122	4.41004452	-0.0078158523	-0.0487268109
## 96	5.08015594	-0.0089454399	-0.0561899263
## 16	11.06966834	-0.0182679640	-0.1235655905
## 39	19.74085320	-0.0313431748	-0.2108471731
## 104	82.54502836	-0.2103654214	-0.3968855837
## 29	16.72887576	-0.0263503782	-0.1861070159
## 71	79.46948218	-0.2114759643	-0.5069152114
## 15	22.48102556	-0.0344777419	-0.2472160284
## 34	25.22455345	-0.0383900372	-0.2755990785
## 119	73.69401660	-0.1862658555	-0.2676293222
## 65	-100.02326809	0.1750895921	0.4623489442
## 91	-82.68369317	0.1732261340	0.5528137735
## 63	-83.77929709	0.1676353390	0.5183505487
## 107	54.52263239	-0.1694459383	-0.5759494216
## 64	25.12954217	-0.0885749956	-0.0467231656
## 67	-39.01050775	0.0592790369	0.2051042654
## 58	35.66599126	-0.1261669497	-0.4989858552
## 120	43.51544091	-0.1379908263	-0.4396728066
## 111	-0.09413534	-0.0322496580	-0.0196023412
## 66	28.70884997	-0.1007363986	-0.3300284050
## 131	-32.11550846	0.0974113483	0.3888868324
## 68	27.41415694	-0.0934513760	-0.1711742602
## 106	-24.81436000	0.0880314038	0.3646556978
## 130	-39.77728309	0.0934919005	0.3498516275
## 98	-22.80619481	0.1091107860	0.4681242005
## 56	-38.65531884	0.1024489672	0.3931534643
## 62	22.69659817	-0.0828540379	-0.2049449464
## 112	-8.65403808	0.0146259551	0.0785762766
## 126	2.07655839	-0.0172354832	-0.0458231645
## 53	-5.68305605	-0.0141200085	0.0152281508
## 115	-27.42134377	0.0608156280	0.2406847141
## 117	13.23178442	-0.0392944275	-0.1912866231
## 33	7.76717018	-0.0170549400	-0.0780114119
## 99	-21.27742723	0.0896608522	0.3764918093
## 108	-17.66314246	0.0667728402	0.2796688503
## 127	-18.84934672	0.0889246482	0.3776454724
## 95	3.90927248	-0.0172890406	-0.0617326941
## 57	18.72368927	-0.0665833840	-0.3478239749
## 102	8.94791098	0.0162173705	0.0663144160
## 100	6.46642748	-0.0176731244	-0.0796954323
## 8	21.00275953	-0.0341884834	-0.2220286045
## 125	0.94423040	0.0616232248	0.2750909498
## 123	14.48365925	-0.0240810860	-0.1504971351
## 14	28.30126764	-0.0507628296	-0.3640891219

## 23	23.57033465	-0.0587978372	-0.4209700874
## 54	26.91834003	0.0335802823	0.1577805812
## 52	-67.67100661	0.1566168121	0.4940275538
## 2	44.47104159	-0.2093412299	-0.9483670010
## 97	17.01145598	0.0695919096	0.3278387503
## 18	-55.19423844	0.0613897925	0.0655661812
## 92	23.10221450	0.0488304742	0.2241374009
## 19	33.70262975	-0.0353936479	-0.2760913646
## 114	43.00613158	-0.0080337664	-0.0770408973
## 101	30.33459374	0.0385482970	0.1727547366
## 25	45.54786275	-0.0181798087	-0.1408979644
## 45	32.77538137	0.0387162952	0.1750980276
## 43	-40.28095395	0.0260437090	-0.0280308221
## 36	-33.98666867	0.0608699393	0.0494886599
## 90	44.54638473	-0.0004402106	-0.0441008572
## 46	42.83192412	-0.0848684592	-0.7062088924
## 61	14.00717860	-0.0904596881	-0.4748269899
## 17	-11.76250713	0.0091272765	-0.1071188917
## 118	22.77491566	-0.0530521030	-0.1488778949
## 60	-56.58679297	0.0456054836	0.1471054576
## 113	-53.68880232	0.1037904471	0.2265014278
## 47	34.02658996	-0.1759375494	-0.4801123988
## 103	83.84105200	-0.1681926199	-0.2269882342
## 30	14.91679728	-0.0871042149	-0.2207281986
## 94	10.29562027	-0.0894073113	-0.4773824082
## 50	5.43907984	-0.1424575183	-0.7240886416
## 32	13.70577437	-0.0169527928	-0.2344326840
## 129	-27.91755348	0.0515999792	0.1417999253
## 51	-28.64853876	-0.0640683984	-0.3678230912
## 31	47.11179378	-0.1285115754	-0.2860008118
## 3	-24.57320923	0.0418710341	-0.0764804713
## 1	-42.03344810	0.0840974327	0.3223150096
## 109	-38.20615560	-0.0101696335	0.2267788759
## 21	-29.86623126	0.0415670775	0.0007227505
## 59	-29.73983420	-0.0209739806	-0.2607241769
## 116	-33.50097865	0.0308018993	0.2419064932
## 48	-5.75087978	-0.0092515444	0.0695525327
## 38	-23.61611474	-0.0130562339	-0.5448368463
## 93	-6.70133976	0.0482401767	0.1395892740
## 41	75.40440608	-0.1352248712	-0.3384178502
## 20	44.15527750	-0.1216279154	-0.5955794570
## 105	-0.05858524	-0.0627155281	-0.6704132068
## 5	49.66620144	-0.0664264716	-0.0413644047
## 4	31.54156751	-0.1671937643	-0.4652021951
## 128	76.73694528	-0.1571835898	-0.1150986143
## 42	4.52341585	0.0229754547	-0.1585003277
## 49	71.10607154	-0.2180382471	-0.5207899642
## 44	33.96846623	-0.0554912227	-0.2307003625
## 22	12.53298864	0.0340155131	0.1034585949
## 13	70.69343157	-0.1646342085	-0.7143871605
## 12	1.89267094	-0.0052933949	0.0090573986
## 9	42.72483895	-0.0391283441	-0.1805153722
## 40	70.96688093	-0.0674434095	-0.1218287269
## 26	44.95547912	-0.1469194627	0.0095931128

```

## 132      -23.69567555  0.0807245820  0.3034811306
## 37       83.52697465 -0.1569631880 -0.5355728353
## 11       44.88516203 -0.0249299264 -0.2639530728
## 10      106.70572989 -0.2289074646 -0.4929933572
## 24       44.28354951 -0.0497220001  0.2503322309

coef(sexmod.l)[1,3]

## [1] 0.6351284

Axes <- seq(100,450,by=50)
Years <- seq(0,10,by=1)

## plot individual fish data
## plot the fixed parameter model
## plot individual fish models

### rgb(0,1,0,0.25, maxColorValue=1)

plot(jitter(LMBL$Agei),LMBL$BI.len,
     col=ifelse(LMBL$Sex==1,rgb(0,0,1,0.25, maxColorValue=1),rgb(1,0,0,0.25, maxColorValue=1)),
     pch=19,
     ylim=c(100,500),
     xlim=c(0,10),
     xlab = "Age (Years)",
     ylab = "Back-Calculated Length (mm)",
     bty="n",
     yaxt="n",
     xaxt="n")

axis(2,at = Axes)
axis(1,at = Years)

abline(h=425.4478748,lty=2,col="blue") ### Males
abline(h=439.0623179,lty=2,col="red") ### females

x <- seq(1,11,by=1)
lines(x, fixef(sexmod.l)[1] * (1 - exp(-fixef(sexmod.l)[3] * (x - (fixef(sexmod.l)[4])))),
      lwd=3,
      col="blue") ### Males

for(i in 1:125){
  lines(x, coef(sexmod.l)[i,1] * (1 - exp(- coef(sexmod.l)[i,3]
    * (x - ( coef(sexmod.l)[i,4] )))),lwd=3,col=rgb(0,0,1,0.1),lty=3) } ### Males

lines(x, fixef(sexmod.l)[2] * (1 - exp(-fixef(sexmod.l)[3] * (x - (fixef(sexmod.l)[4])))),
      lwd=3,
      col="red") ### Females

for(i in 1:125){
  lines(x, coef(sexmod.l)[i,2] * (1 - exp(- coef(sexmod.l)[i,3]
    * (x - ( coef(sexmod.l)[i,4] )))),lwd=3,col=rgb(1,0,0,0.1),lty=3) } ### Females

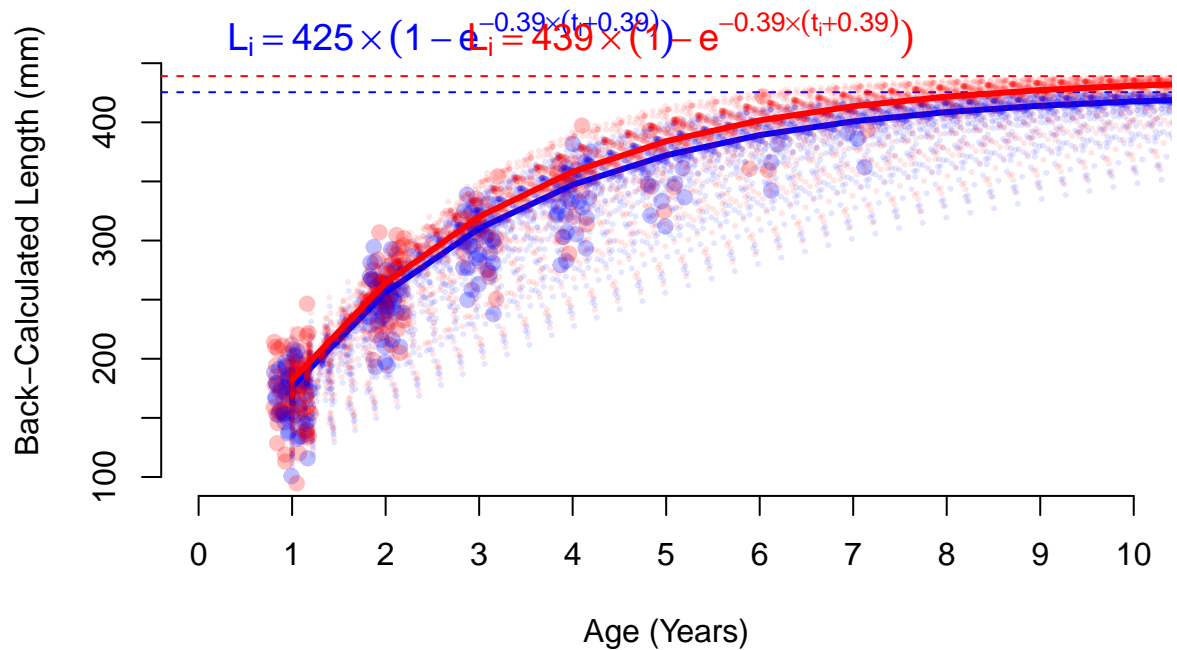
legend("topleft",
      legend = print(expression(L[i]==425 %*% (1 - e **{-0.39 %*% (t[i] + 0.39)}))),

```

```
bty="n",
cex=1.15,
text.col = "blue") ### VB Equation Males
```

```
## expression(L[i] == 425 %*% (1 - e^{
##   -0.39 %*% (t[i] + 0.39)
## })))
```

```
legend("top",
  legend = print(expression(L[i]==439 %*% (1 - e **{-0.39 %*% (t[i] + 0.39)}))),
  bty="n",
  cex=1.15,
  text.col = "red") ### VB Equation Females
```



```
## expression(L[i] == 439 %*% (1 - e^{
##   -0.39 %*% (t[i] + 0.39)
## })))
```

```
{sexmod.l.int}
```

```
## [1] "Iterations = 8"
```

```
## Nonlinear mixed-effects model fit by REML
```

```
## Model: BL.len ~ LVB(Agei, Linf, K, t0)
```

```
## Data: datgr
```

```
## AIC BIC logLik
```

```
## 2809.756 2851.613 -1393.878
```

```
##
```

```
## Random effects:
```

```
## Formula: list(Linf ~ 1, K ~ 1, t0 ~ 1)
```

```
## Level: FID
```

```
## Structure: General positive-definite, Log-Cholesky parametrization
```

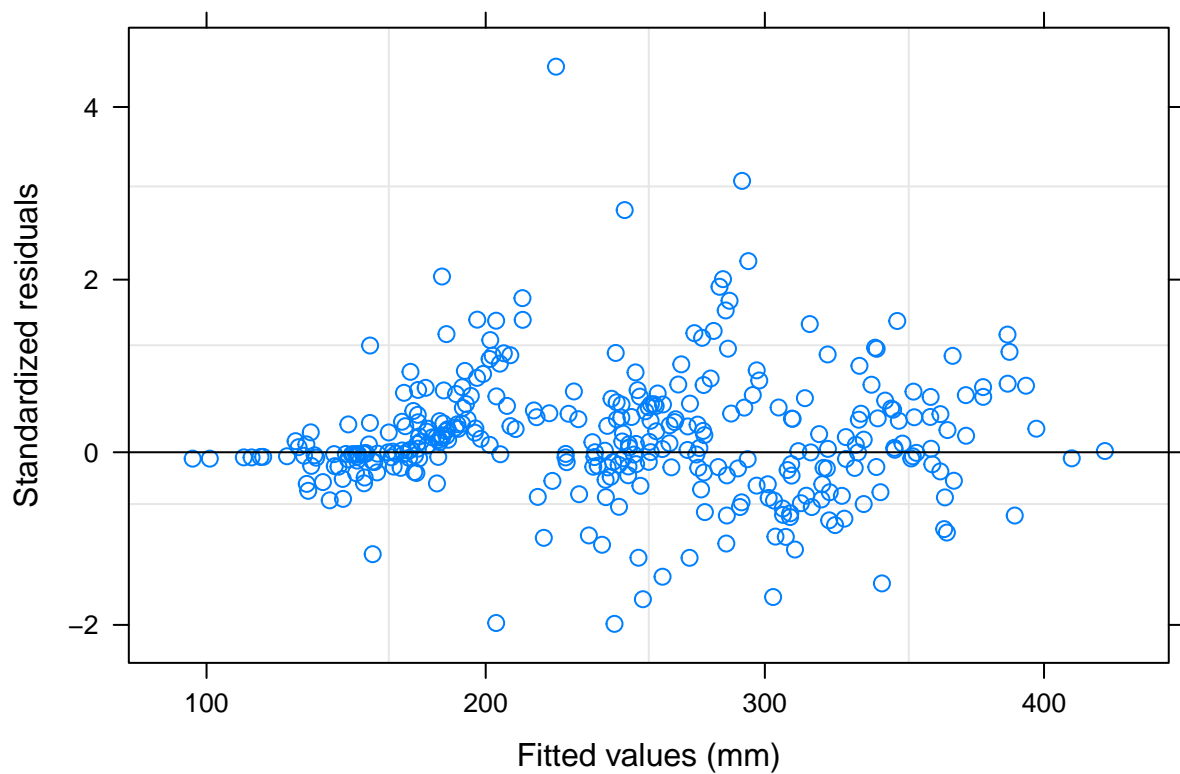
```
## StdDev Corr
```

```
## Linf.(Intercept) 60.7967245 Ln.(I) K
```

```

## K                0.1283638 -0.880
## t0               0.4334913 -0.675  0.857
## Residual        4.7920262
##
## Fixed effects: list(Linf ~ Sex, K ~ 1, t0 ~ 1)
##               Value Std.Error DF   t-value p-value
## Linf.(Intercept) 425.4479  8.432969 208 50.45055  0.0000
## Linf.Sex2        13.6145  6.626469 208  2.05456  0.0412
## K                0.3855  0.015426 208 24.98990  0.0000
## t0              -0.3887  0.043652 208 -8.90441  0.0000
## Correlation:
##      Ln.(I) Lnf.S2 K
## Linf.Sex2 -0.376
## K         -0.824  0.004
## t0        -0.582  0.040  0.811
##
## Standardized Within-Group Residuals:
##      Min      Q1      Med      Q3      Max
## -1.9880697 -0.1671674  0.0861218  0.4965188  4.4670194
##
## Number of Observations: 336
## Number of Groups: 125
##
## Approximate 95% confidence intervals
##
## Fixed effects:
##               lower      est.      upper
## Linf.(Intercept) 408.8228916 425.4479383 442.0729850
## Linf.Sex2        0.5508558  13.6145061  26.6781564
## K                0.3550873  0.3854991  0.4159108
## t0              -0.4747486 -0.3886923 -0.3026359
## attr("label")
## [1] "Fixed effects:"
##
## Random Effects:
## Level: FID
##               lower      est.      upper
## sd(Linf.(Intercept)) 47.2998329 60.7967245 78.1449211
## sd(K)                0.0822892  0.1283638  0.2002362
## sd(t0)               0.3685501  0.4334913  0.5098757
## cor(Linf.(Intercept),K) -0.9538721 -0.8799822 -0.7055677
## cor(Linf.(Intercept),t0) -0.8543408 -0.6745775 -0.3506498
## cor(K,t0)            0.7122032  0.8565806  0.9314297
##
## Within-group standard error:
##      lower      est.      upper
## 3.979148 4.792026 5.770963
##
##               numDF denDF   F-value p-value
## Linf.(Intercept)    1   208 41958.70 <.0001
## Linf.Sex            1   208   25.59 <.0001
## K                   1   208 3026.00 <.0001
## t0                  1   208   79.29 <.0001

```



```
fixef(sexmod.l.int)
```

##	Linf.(Intercept)	Linf.Sex2	K	t0
##	425.4479383	13.6145061	0.3854991	-0.3886923

```
ranef(sexmod.l.int)
```

##	Linf.(Intercept)	K	t0
## 88	-96.09321168	0.249629216	0.8778497795
## 87	-89.71646144	0.229926649	0.8060792709
## 85	-82.95977400	0.211558879	0.7468872657
## 86	-77.70719271	0.195853142	0.6897164842
## 83	-78.14581280	0.197982078	0.7007738464
## 27	-77.33190265	0.195704451	0.6930648794
## 69	-69.82374377	0.174919577	0.6230374893
## 84	-64.00491304	0.159070714	0.5699789812
## 82	-55.36027694	0.135284052	0.4867700371
## 110	-41.59316212	0.099405607	0.3675619827
## 76	-44.97302255	0.108590018	0.4020757671
## 80	-44.23795275	0.106678158	0.3957217311
## 74	-43.08489280	0.103684495	0.3857686046
## 78	-42.36037907	0.101806889	0.3795229856
## 77	-42.07766547	0.101074941	0.3770876183
## 75	-39.61724203	0.094721427	0.3559281313
## 72	-31.44864984	0.073585352	0.2812061590
## 121	-38.36871533	0.091508864	0.3452125665
## 73	-14.78547244	0.032645824	0.1416830699
## 7	-9.33238402	0.019907888	0.0919717676
## 70	2.59650714	-0.004651973	-0.0287298236
## 35	9.05294684	-0.015910237	-0.0958422729

## 122	4.41006677	-0.007815898	-0.0487266969
## 96	5.08018809	-0.008945504	-0.0561898641
## 16	11.06978320	-0.018268166	-0.1235659235
## 39	19.74105967	-0.031343472	-0.2108479542
## 104	82.54496875	-0.210365424	-0.3968868192
## 29	16.72905323	-0.026350657	-0.1861076140
## 71	79.46950468	-0.211476009	-0.5069162518
## 15	22.48124838	-0.034478045	-0.2472167359
## 34	25.22479098	-0.038390341	-0.2755997795
## 119	73.69391981	-0.186265869	-0.2676302733
## 65	-100.02319355	0.175089167	0.4623491570
## 91	-82.68368794	0.173225942	0.5528140347
## 63	-83.77927882	0.167635108	0.5183507967
## 107	54.52279514	-0.169446065	-0.5759499153
## 64	25.12919521	-0.088574597	-0.0467223168
## 67	-39.01039355	0.059278761	0.2051042226
## 58	35.66620325	-0.126167131	-0.4989862363
## 120	43.51558507	-0.137990994	-0.4396733347
## 111	-0.09425749	-0.032249427	-0.0196017741
## 66	28.70896021	-0.100736514	-0.3300286636
## 131	-32.11545849	0.097411401	0.3888871447
## 68	27.41406913	-0.093451318	-0.1711741220
## 106	-24.81437527	0.088031647	0.3646562165
## 130	-39.77736091	0.093491966	0.3498519420
## 98	-22.80627003	0.109110987	0.4681244493
## 56	-38.65539609	0.102449091	0.3931538749
## 62	22.69661412	-0.082854113	-0.2049450236
## 112	-8.65412241	0.014626145	0.0785767240
## 126	2.07651641	-0.017235379	-0.0458228154
## 53	-5.68314390	-0.014119918	0.0152285630
## 115	-27.42143945	0.060815746	0.2406850924
## 117	13.23188265	-0.039294551	-0.1912867296
## 33	7.76716359	-0.017054895	-0.0780112098
## 99	-21.27755277	0.089661157	0.3764923833
## 108	-17.66328388	0.066773074	0.2796692712
## 127	-18.84954620	0.088925050	0.3776460809
## 95	3.90924046	-0.017289038	-0.0617324750
## 57	18.72393853	-0.066583766	-0.3478245719
## 102	8.94783747	0.016217557	0.0663146882
## 100	6.46641009	-0.017673142	-0.0796952723
## 8	21.00286562	-0.034188583	-0.2220287689
## 125	0.94408297	0.061623499	0.2750912683
## 123	14.48369588	-0.024081171	-0.1504971693
## 14	28.30144864	-0.050762933	-0.3640893678
## 23	23.57054539	-0.058798029	-0.4209703548
## 54	26.91830145	0.033580509	0.1577806102
## 52	-67.67100387	0.156616739	0.4940278079
## 2	44.47128270	-0.209341310	-0.9483672770
## 97	17.01134313	0.069592204	0.3278389153
## 18	-55.19409047	0.061389300	0.0655657506
## 92	23.10212166	0.048830737	0.2241375045
## 19	33.70270698	-0.035393685	-0.2760915941
## 114	43.00613990	-0.008033666	-0.0770411406
## 101	30.33452595	0.038548498	0.1727546169


```
## 25      45.54789055 -0.018179758 -0.1408982465
## 45      32.77531900  0.038716509  0.1750978762
## 43     -40.28076273  0.026043041 -0.0280316648
## 36     -33.98660907  0.060869745  0.0494886303
## 90      44.54633812 -0.000440037 -0.0441009923
## 46      42.83279728 -0.084869076 -0.7062110233
## 61      14.00756590 -0.090460297 -0.4748284428
## 17     -11.76237532  0.009126989 -0.1071192445
## 118     22.77487750 -0.053052020 -0.1488776381
## 60     -56.58692157  0.045605649  0.1471060589
## 113    -53.68883477  0.103790475  0.2265017561
## 47      34.02670528 -0.175937613 -0.4801129254
## 103     83.84098477 -0.168192527 -0.2269882642
## 30      14.91681887 -0.087104238 -0.2207282512
## 94      10.29576932 -0.089407526 -0.4773829926
## 50       5.43918603 -0.142457607 -0.7240889652
## 32      13.70590016 -0.016953019 -0.2344331770
## 129    -27.91761112  0.051600105  0.1418003821
## 51     -28.64853320 -0.064068421 -0.3678230268
## 31      47.11181097 -0.128511634 -0.2860011019
## 3      -24.57322697  0.041871051 -0.0764802187
## 1      -42.03358631  0.084097628  0.3223155713
## 109    -38.20639930 -0.010169173  0.2267800448
## 21     -29.86633252  0.041567161  0.0007231562
## 59     -29.73986759 -0.020973920 -0.2607238433
## 116    -33.50107675  0.030802150  0.2419071911
## 48      -5.75106398 -0.009251272  0.0695532693
## 38     -23.61608258 -0.013056306 -0.5448368653
## 93      -6.70140794  0.048240332  0.1395897464
## 41      75.40449262 -0.135224941 -0.3384182792
## 20      44.15542456 -0.121628143 -0.5955802976
## 105    -0.05855515 -0.062715513 -0.6704130066
## 5      49.66613827 -0.066426352 -0.0413641175
## 4      31.54159520 -0.167193820 -0.4652027309
## 128     76.73685725 -0.157183462 -0.1150984113
## 42       4.52340652  0.022975477 -0.1585001555
## 49      71.10611590 -0.218038213 -0.5207903342
## 44      33.96849902 -0.055491259 -0.2307004581
## 22      12.53292218  0.034015664  0.1034590390
## 13      70.69356966 -0.164634298 -0.7143878331
## 12       1.89260155 -0.005293252  0.0090578443
## 9       42.72486870 -0.039128366 -0.1805154458
## 40      70.96693576 -0.067443468 -0.1218289491
## 26      44.95539192 -0.146919319  0.0095934612
## 132    -23.69573821  0.080724696  0.3034814946
## 37      83.52707432 -0.156963302 -0.5355736071
## 11      44.88516133 -0.024929947 -0.2639532188
## 10     106.70574808 -0.228907441 -0.4929939169
## 24      44.28339194 -0.049721817  0.2503327326
```

```
coef(sexmod.l.int)[1,3]
```

```
## [1] 0.6351283
```

Comparing Models

```
AIC(nlme.mod2,sexmod.lt,sexmod.kt,sexmod.l,sexmod.k,sexmod.t,sexmod.l.int)
```

```
## Warning in AIC.default(nlme.mod2, sexmod.lt, sexmod.kt, sexmod.l,  
## sexmod.k, : models are not all fitted to the same number of observations
```

```
##           df      AIC  
## nlme.mod2    10 2825.199  
## sexmod.lt    12 2827.788  
## sexmod.kt    12 2837.204  
## sexmod.l     11 2809.756  
## sexmod.k     11 2824.301  
## sexmod.t     11 2829.548  
## sexmod.l.int 11 2809.756
```

```
BIC(nlme.mod2,sexmod.lt,sexmod.kt,sexmod.l,sexmod.k,sexmod.t,sexmod.l.int)
```

```
## Warning in BIC.default(nlme.mod2, sexmod.lt, sexmod.kt, sexmod.l,  
## sexmod.k, : models are not all fitted to the same number of observations
```

```
##           df      BIC  
## nlme.mod2    10 2863.280  
## sexmod.lt    12 2873.413  
## sexmod.kt    12 2882.830  
## sexmod.l     11 2851.613  
## sexmod.k     11 2866.158  
## sexmod.t     11 2871.404  
## sexmod.l.int 11 2851.613
```

```
anova(nlme.mod2,sexmod.lt,sexmod.kt,sexmod.l,sexmod.k,sexmod.t,sexmod.l.int)
```

```
##           Model df      AIC      BIC    logLik    Test    L.Ratio p-value  
## nlme.mod2      1 10 2825.199 2863.280 -1402.599  
## sexmod.lt      2 12 2827.788 2873.413 -1401.894 1 vs 2  1.410824  0.4939  
## sexmod.kt      3 12 2837.204 2882.830 -1406.602  
## sexmod.l       4 11 2809.756 2851.613 -1393.878 3 vs 4 25.448099 <.0001  
## sexmod.k       5 11 2824.301 2866.157 -1401.150  
## sexmod.t       6 11 2829.548 2871.404 -1403.774  
## sexmod.l.int   7 11 2809.756 2851.613 -1393.878
```

```
anova(nlme.mod2,sexmod.l,sexmod.l.int)
```

```
##           Model df      AIC      BIC    logLik    Test    L.Ratio p-value  
## nlme.mod2      1 10 2825.199 2863.280 -1402.599  
## sexmod.l       2 11 2809.756 2851.613 -1393.878 1 vs 2 17.4423 <.0001  
## sexmod.l.int   3 11 2809.756 2851.613 -1393.878
```

```
anova(sexmod.l,nlme.mod2,sexmod.l.int)
```

```
##           Model df      AIC      BIC    logLik    Test    L.Ratio p-value  
## sexmod.l       1 11 2809.756 2851.613 -1393.878  
## nlme.mod2      2 10 2825.199 2863.280 -1402.599 1 vs 2 17.44230 <.0001  
## sexmod.l.int   3 11 2809.756 2851.613 -1393.878 2 vs 3 17.44229 <.0001
```

```
anova(sexmod.l,sexmod.k,nlme.mod2)
```

```
##           Model df      AIC      BIC    logLik    Test    L.Ratio p-value  
## sexmod.l       1 11 2809.756 2851.613 -1393.878
```

```
## sexmod.k      2 11 2824.301 2866.157 -1401.150
## nlme.mod2     3 10 2825.199 2863.280 -1402.599 2 vs 3 2.89759 0.0887
```

```
anova(sexmod.l,nlme.mod2)
```

```
##           Model df      AIC      BIC    logLik  Test L.Ratio p-value
## sexmod.l      1 11 2809.756 2851.613 -1393.878
## nlme.mod2     2 10 2825.199 2863.280 -1402.599 1 vs 2 17.4423 <.0001
```

```
anova(nlme.mod2,sexmod.l)
```

```
##           Model df      AIC      BIC    logLik  Test L.Ratio p-value
## nlme.mod2     1 10 2825.199 2863.280 -1402.599
## sexmod.l      2 11 2809.756 2851.613 -1393.878 1 vs 2 17.4423 <.0001
```

```
anova(sexmod.l,sexmod.l.int)
```

```
##           Model df      AIC      BIC    logLik
## sexmod.l      1 11 2809.756 2851.613 -1393.878
## sexmod.l.int  2 11 2809.756 2851.613 -1393.878
```

```
anova(sexmod.l.int,sexmod.l)
```

```
##           Model df      AIC      BIC    logLik
## sexmod.l.int  1 11 2809.756 2851.613 -1393.878
## sexmod.l      2 11 2809.756 2851.613 -1393.878
```

See sexmod.l

There is a significant difference between the nlme.mod2 and sexmod.l with sexmod.l being the model that best describes the data (loglikelihood,df=11, L Ratio = 17.4423, $p < 0.0001$).